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V 0 V
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The list of best scores is:
                                            A 100% identical sequence to the
                                                                       The scores below are sorted by initial score. Significance is calculated based on initial score.
                                                                                                                                   Number of residues:
Number of sequences
Number of scores ab
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Number of sequences searched:
1
Number of scores above cutoff:
                                                                                                                                                                                                                   Times:
                                                                                                                                                                                                                                                                 Scores:
                                                                                                                                                                                                                                                                                                                                        Cur-- C'score
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                                                                                                                                                                                                                                                                                                                        Randomization group
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                                            query sequence was not found
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Joining penalty
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1. US-09-831-805A-20 (1-917)
y14736 TOIG of: y14736 check: 615 from: 1
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AUTHORS
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ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AUTHORS
TITLE
JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JOURNAL
MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (03-SEP-1997) T. Paterson, National Science Laboratory, Scottish Nat. Blood Transfusion Service, 12 Bristo Place, Edinburgh, EH1 1EZ, Scotland, UK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Paterson,T., Innes,J., McMillan,L., Downing,I. and Carter,M.C. Variation in IgGl heavy chain allotype does not contribute to differences in biological activity of two human anti-Rhesus (D)
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Homo sapiens mRNA for immunoglobulin kappa light chain.
Y14736
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                     237
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                 402..725
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region"
274 c 218 g 209 t
                                                                                                                          region
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csD-1_(THERAD 03)"
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840 850 860 TCATCTTCACCTCACCCCCCTCCT	760 770 780 790 CCTCAGTTCCAGCCTGACCCCTCCCATCCTTTGGCCTCTG	690 700 710 AGCTCGCCCGTCACAAGAGCTTCAACAGGGGAG	20 630 640 BACGCTGAGCAAAGCAGACTACGAGAAAC	550 560 570 \$10,000 \$1	480 490 500 RETTGTGTGCCTGCTGAATAACTTCTATC	410 420 420 ACGAACTGTGGCTGCACCATCTGTCTTCA	340 350 GCAACTTACTACTGTCAACAACTTACAGTA	50 270 280 AAGGTTCAGTGGCAGTGGATCTGGGACAG	190 210 TATCAGCAGAAACCAGGGAAAGCCCCTAAGC	120 130 140 1GCATCTGTAGGAGACACAGTCACCATCA 111 1111 11 11 11 11 11 11 11 11 11 11	ဂ္ဂ 🗕 ဂ္ဂ	87 88 tio

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